Please replace the first full paragraph on page 25 with the following amended paragraph:

FIG. 4A, FIG. 4B, FIG. 4C and FIG. 4 FIG. 4D. Active site of ICL. FIG. 4A

and FIG. 4B, stereo view of the $2|F_0|-|F_c|\phi_c$ electron density map contoured at 1σ for bound 3-

bromopyruvate modified ICL. Shown is the active site with the pyruvyl moiety (purple) attached

to the thiolate of Cys191. FIG 4C and FIG. 4 FIG. 4D, stereo view of active site of ICL shown

as a molecular surface colored according to the electrostatic surface potential. The surface was

generated based on the protein coordinates of the ternary complex of ICL with glyoxylate (green)

and succinate (cyan). The atoms of the active site loop (residues 183-197) and the C-terminal

segment (residues 410-427) of the adjacent subunit were excluded during surface calculations.

The loop segments are shown as yellow ribbon for the inhibitor complex and white ribbon for the

apo enzyme. Side-chains of some of the residues have been omitted for clarity, as is standard in

this art.